

Figure 1

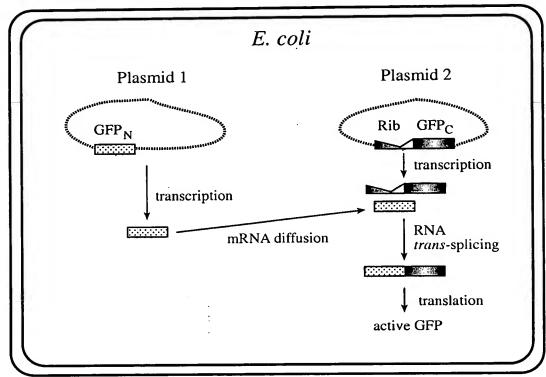


Figure 2

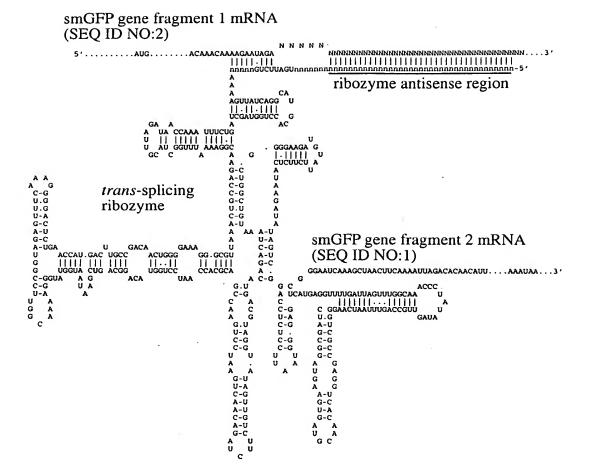
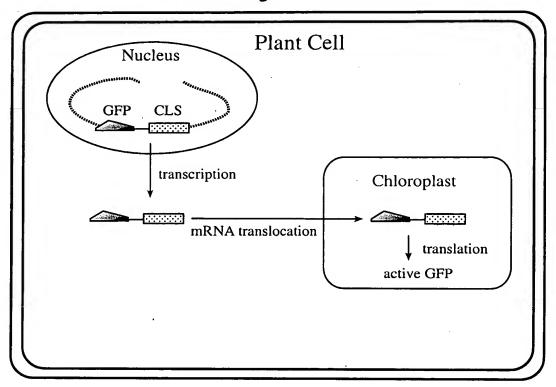


Figure 3

Figure 4



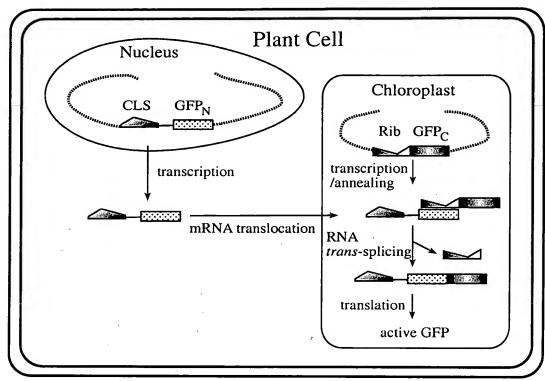


Figure 5

Figure 6

SEQ ID NO:3

tttattagaa caagaagtga ggatatgatt aaactttgtt tgacgaaacc
aggtctgttc cgactttccg actctgagtt tcgacttgtg agagaaggag
gagtcgtggt gaacttttat taaaaaaatt agttcactcg tcttcaatct
cttgatcact tcgtctctc agggaaagat gggaagaaca ctgatgagtc

201 tcgcaaggtt tactcctcta tcttcattgt ttttttacaa aatcttg

Figure 7

5'- LB ASBVd(+) UTR smGFP RB -3'

Figure 8A

pVUG1 seqment 5689-6757 (SEQ ID NO:4)

5689	tttattagaa	caagaagtga	ggatatgatt	aaactttgtt	tgacgaaacc
5739	aggtctgttc	cgactttccg	actctgagtt	tcgacttgtg	agagaaggag
5789	gagtcgtggt	gaacttttat	taaaaaaatt	agttcactcg	tcttcaatct
5839	cttgatcact	tcgtgctagc	atgtatttgg	caaatcaaat	accatggtct
5889	aataatcaaa	cattctgatt	agttgataat	attagtatta	gttggaaatt
5939	ttgtgaaaga	ttcctatgaa	aagtttcatt	aacacggaat	tcgtgtcgag
5989	tagaccttgt	tgttgtgaga	attcttaatt	catgagttgt	agggagggat
6039	ttATGAGTAA	AGGAGAAGAA	CTTTTCACTG	GAGTTGTCCC	AATTCTTGTT
6089	GAATTAGATG	GTGATGTTAA	TGGGCACAAA	TTTTCTGTCA	GTGGAGAGGG
6139	TGAAGGTGAT	GCAACATACG	GAAAACTTAC	CCTTAAATTT	ATTTGCACTA
6189	CTGGAAAACT	ACCTGTTCCA	TGGCCAACAC	TTGTCACTAC	TTTCTCTTAT
6239	GGTGTTCAAT	GCTTTTCAAG	ATACCCAGAT	CATATGAAGC	GGCACGACTT
6289	CTTCAAGAGC	GCCATGCCTG	AGGGATACGT	GCAGGAGAGG	ACCATCTCTT
6339	TCAAGGACGA	CGGGAACTAC	AAGACACGTG	CTGAAGTCAA	GTTTGAGGGA
6389	GACACCCTCG	TCAACAGGAT	CGAGCTTAAG	GGAATCGATT	TCAAGGAGGA
6439	CGGAAACATC	CTCGGCCACA	AGTTGGAATA	CAACTACAAC	TCCCACAACG
6489	TATACATCAC	GGCAGACAAA	CAAAAGAATG	GAATCAAAGC	TAACTTCAAA
6539	ATTAGACACA	ACATTGAAGA	TGGAAGCGTT	CAACTAGCAG	ACCATTATCA
6589	ACAAAATACT	CCAATTGGCG	ATGGCCCTGT	CCTTTTACCA	GACAACCATT
6639	ACCTGTCCAC	ACAATCTGCC	CTTTCGAAAG	ATCCCAACGA	AAAGAGAGAC
6689	CACATGGTCC	TTCTTGAGTT	TGTAACAGCT	GCTGGGATTA	CACATGGCAT
6739	GGATGAACTA	ТАСАААТАА			

Figure 8B

pVUG2 seqment 5689-6757 (SEQ ID NO:5)

5689	acttgtgaga	gaaggaggag	tcgtggtgaa	cttttattaa	aaaaattagt
5739	tcactcgtct	tcaatctctt	gatcacttcg	tctcttcagg	gaaagatggg
5789	aagaacactg	atgagtctcg	caaggtttac	tcctctatct	tcattgtttt
5839	tttacaaaat	cttggctagc	atgtatttgg	caaatcaaat	accatggtct
5889	aataatcaaa	cattctgatt	agttgataat	attagtatta	gttggaaatt
5939	ttgtgaaaga	ttcctatgaa	aagtttcatt	aacacggaat	tcgtgtcgag
5989	tagaccttgt	tgttgtgaga	attcttaatt	catgagttgt	agggagggat
6039	ttATGAGTAA	AGGAGAAGAA	CTTTTCACTG	GAGTTGTCCC	AATTCTTGTT
6089	GAATTAGATG	GTGATGTTAA	TGGGCACAAA	TTTTCTGTCA	GTGGAGAGGG
6139	TGAAGGTGAT	GCAACATACG	GAAAACTTAC	ССТТАААТТТ	ATTTGCACTA
6189	CTGGAAAACT	ACCTGTTCCA	TGGCCAACAC	TTGTCACTAC	TTTCTCTTAT
6239	GGTGTTCAAT	GCTTTTCAAG	ATACCCAGAT	CATATGAAGC	GGCACGACTT
6289	CTTCAAGAGC	GCCATGCCTG	AGGGATACGT	GCAGGAGAGG	ACCATCTCTT
6339	TCAAGGACGA	CGGGAACTAC	AAGACACGTG	CTGAAGTCAA	GTTTGAGGGA
6389	GACACCCTCG	TCAACAGGAT	CGAGCTTAAG	GGAATCGATT	TCAAGGAGGA
6439	CGGAAACATC	CTCGGCCACA	AGTTGGAATA	CAACTACAAC	TCCCACAACG
6489	TATACATCAC	GGCAGACAAA	CAAAAGAATG	GAATCAAAGC	TAACTTCAAA
6539	ATTAGACACA	ACATTGAAGA	TGGAAGCGTT	CAACTAGCAG	ACCATTATCA
6589	АСААААТАСТ	CCAATTGGCG	ATGGCCCTGT	CCTTTTACCA	GACAACCATT
6639	ACCTGTCCAC	ACAATCTGCC	CTTTCGAAAG	ATCCCAACGA	AAAGAGAGAC
6689	CACATGGTCC	TTCTTGAGTT	TGTAACAGCT	GCTGGGATTA	CACATGGCAT
6739	GGATGAACTA	ТАСАААТАА			

Figure 8C

pVUG3 segment 6833-7901 (SEQ ID NO:6)

6833	tcttcaggga	aagatgggaa	gaacactgat	gagtctcgca	aggtttactc
6883	ctctatcttc	attgttttt	tacaaaatct	tgtttattag	aacaagaagt
6933	gaggatatga	ttaaactttg	tttgacgaaa	ccaggtctgt	tccgactttc
6983	cgactctgag	tttcgctagc	atgtatttgg	caaatcaaat	accatggtct
7033	aataatcaaa	cattctgatt	agttgataat	attagtatta	gttggaaatt
7083	ttgtgaaaga	ttcctatgaa	aagtttcatt	aacacggaat	tcgtgtcgag
7133	tagaccttgt	tgttgtgaga	attcttaatt	catgagttgt	agggagggat
7183	ttATGAGTAA	AGGAGAAGAA	CTTTTCACTG	GAGTTGTCCC	AATTCTTGTT
7233	GAATTAGATG	GTGATGTTAA	TGGGCACAAA	TTTTCTGTCA	GTGGAGAGGG
7283	TGAAGGTGAT	GCAACATACG	GAAAACTTAC	CCTTAAATTT	ATTTGCACTA
7333	CTGGAAAACT	ACCTGTTCCA	TGGCCAACAC	TTGTCACTAC	TTTCTCTTAT
7383	GGTGTTCAAT	GCTTTTCAAG	ATACCCAGAT	CATATGAAGC	GGCACGACTT
7433	CTTCAAGAGC	GCCATGCCTG	AGGGATACGT	GCAGGAGAGG	ACCATCTCTT
7483	TCAAGGACGA	CGGGAACTAC	AAGACACGTG	CTGAAGTCAA	GTTTGAGGGA
7533	GACACCCTCG	TCAACAGGAT	CGAGCTTAAG	GGAATCGATT	TCAAGGAGGA
7583	CGGAAACATC	CTCGGCCACA	AGTTGGAATA	CAACTACAAC	TCCCACAACG
7633	TATACATCAC	GGCAGACAAA	CAAAAGAATG	GAATCAAAGC	ТААСТТСААА
7683	ATTAGACACA	ACATTGAAGA	TGGAAGCGTT	CAACTAGCAG	ACCATTATCA
7733	ACAAAATACT	CCAATTGGCG	ATGGCCCTGT	CCTTTTACCA	GACAACCATT
7783	ACCTGTCCAC	ACAATCTGCC	CTTTCGAAAG	ATCCCAACGA	AAAGAGAGAC
7833	CACATGGTCC	TTCTTGAGTT	TGTAACAGCT	GCTGGGATTA	CACATGGCAT
7883	GGATGAACTA	TACAAATAA			

Figure 9



Western blot analysis using anti-GFP antibody

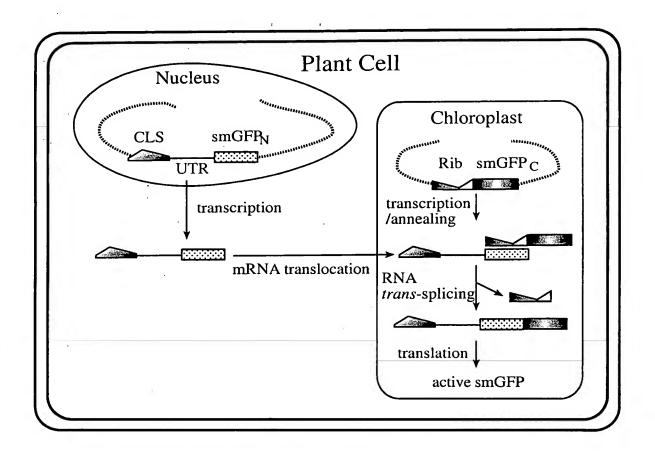


Figure 10

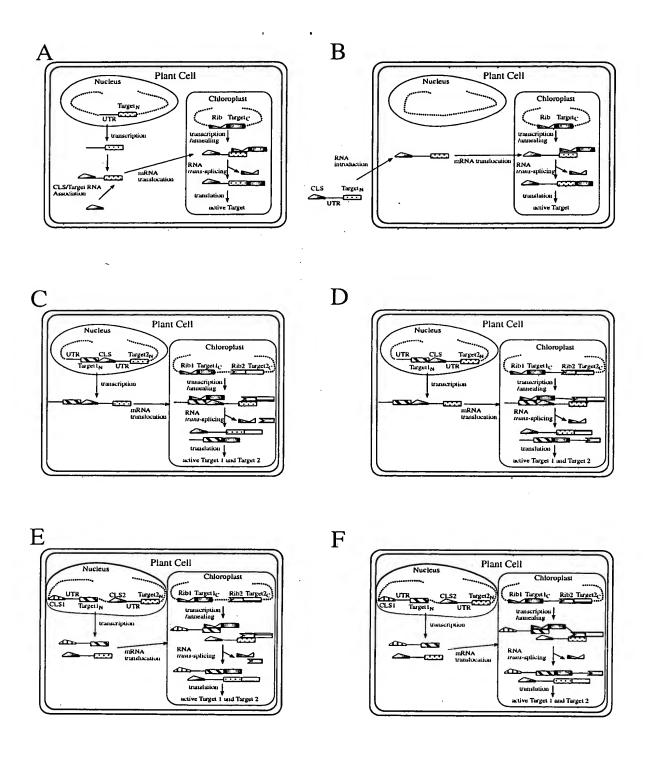


Figure 11